

FIGURE 1

10007075-422000

AGamma_1:

hinge Ch₂

Native A E P K S C D K T H T C P P C P..... A P E L L G G P S.....
 234 235 237

Mutated A E P K S S D K T H T S P P S P..... A P E A E G A P S.....
 234 235 237

3Gamma_4:

hinge Ch₂

Native A E S K Y G P P C P S C P..... A P E F L G G P S.....
 234 235 237

Mutated A E S K Y G P P S P S S P..... A P E F G G A P S.....
 234 235 237

FIGURE 2

100227075 - 100

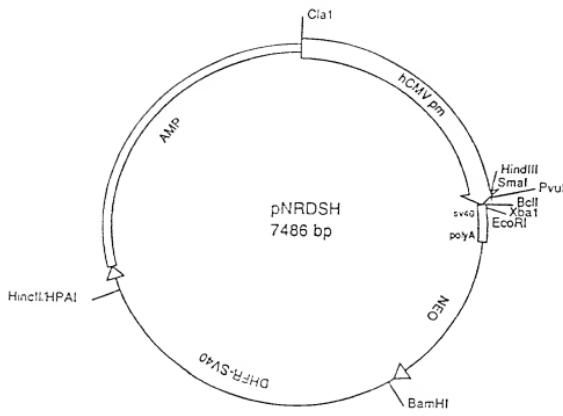
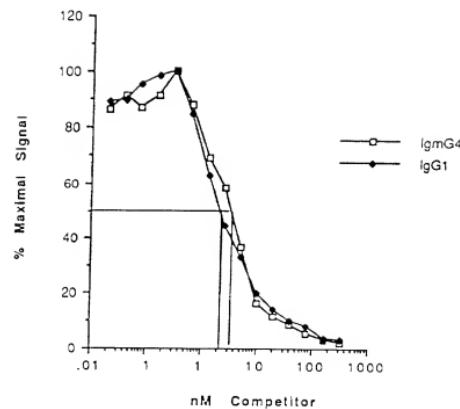


FIGURE 3

20027075-422004

A.



B.

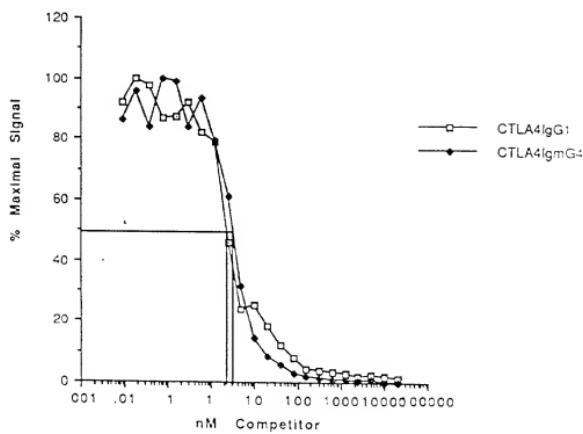
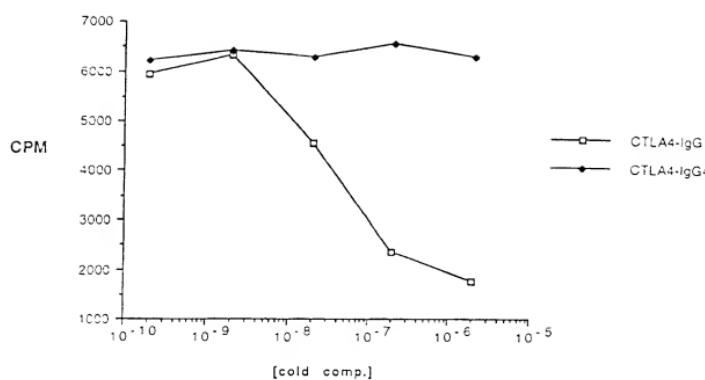


FIGURE 4

FIGURE 5

A



B

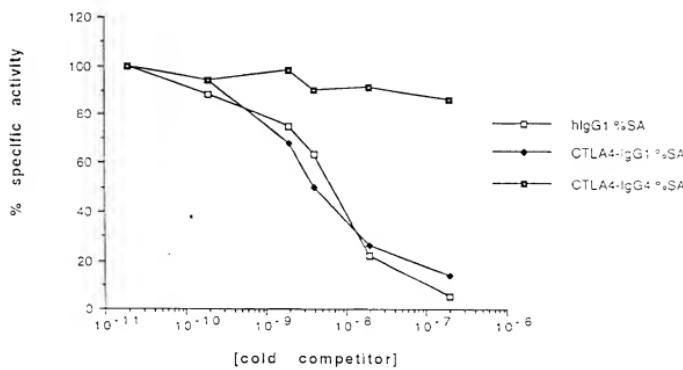


FIGURE 5

Y00227-570200Y

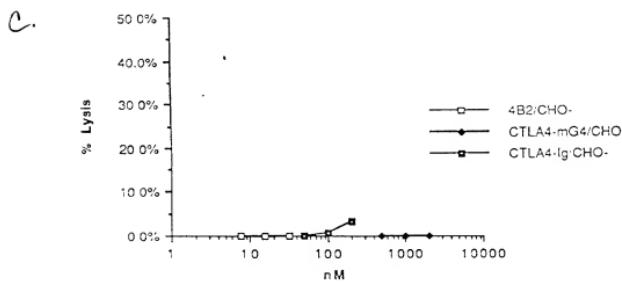
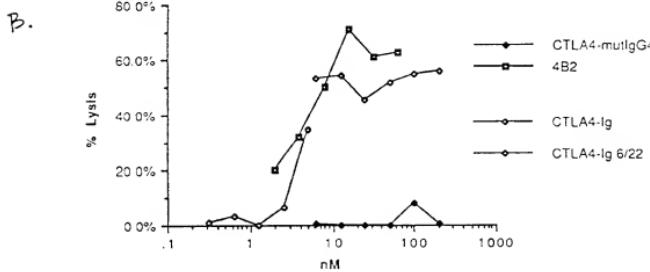
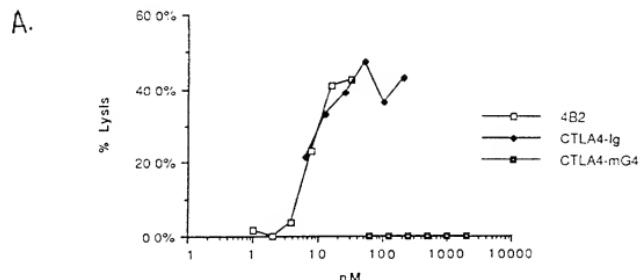


FIGURE 6

10007075-1223001

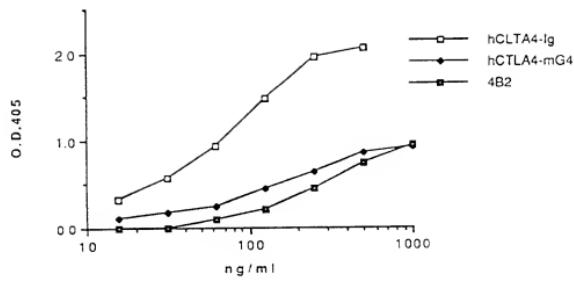


FIGURE 7

YODZER 5/20/2007

FIGURE 8

